IN THE UNITED STATES PATENT AND TRADEMARK OFFICE
(Case No. 00,1214)

PATENT

In re Application of:	**Welcher et al.**	)
		)
Serial No.:	**Unassigned**	) **Before the Examiner:** **Unassigned**
		)
Filed:	**November 28, 2000**	) **Group Art Unit:** **Unassigned**
		)
For:	**Interleukin-1**	)
	Receptor Antagonist-Like	)
	Molecules and Uses Thereof	)

Assistant Commissioner for Patents
Washington, D.C. 20231

Sir/Madam:

STATEMENT UNDER 37 C.F.R. § 1.821

The undersigned representative hereby declares that the content of the paper and computer readable copies of the Sequence Listing, submitted in the above-identified application in accordance with 37 C.F.R. §§ 1.821(c) and (e), respectively, are identical in content. The 3.5" diskette contains an IBM compatible dos-text file of the sequence listing named "001214seq.txt."

Respectfully submitted,
McDonnell Boehnen Hulbert & Berghoff

Dated: November 28, 2000

By: _____
Kevin E. Noonan, Ph.D.
Reg. No. 35,303

SEQUENCE LISTING

<110> Welcher, Andrew A.
Jing, Shuqian
Luethy, Roland

<120> Interleukin-1 Receptor Antagonist-Like Molecules and Uses Thereof

<130> 00-1214

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<150> 60/170,052
<151> 1999-12-10

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<170> PatentIn Ver. 2.0

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cagcctcctc accaccatct gatctatctt gttctcttca caaaaggctc tgaagacatc 300

atg aac cca caa cgg gag gca gca ccc aaa tcc tat gct att cgt gat 348
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1 5 10 15

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Ser Arg Gln Met Val Trp Val Leu Ser Gly Asn Ser Leu Ile Ala Ala
20 25 30

cct ctt agc cgc agc att aag cct gtc act ctt cat tta ata gcc tgt 444
Pro Leu Ser Arg Ser Ile Lys Pro Val Thr Leu His Leu Ile Ala Cys
35 40 45

aga gac aca gaa ttc agt gac aag gaa aag ggt aat atg gtt tac ctg 492
Arg Asp Thr Glu Phe Ser Asp Lys Glu Lys Gly Asn Met Val Tyr Leu
50 55 60

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aag cct act ttg cag ctt aag gaa aaa aat atc atg gac ctg tat gtg   588
Lys Pro Thr Leu Gln Leu Lys Glu Lys Asn Ile Met Asp Leu Tyr Val
                 85                  90                  95

gag aag aaa gca cag aag ccc ttt ctc ttt ttc cac aat aaa gaa ggc   636
Glu Lys Lys Ala Gln Lys Pro Phe Leu Phe Phe His Asn Lys Glu Gly
            100                 105                 110

tcc act tct gtc ttt cag tca gtc tct tac cct ggc tgg ttc ata gcc   684
Ser Thr Ser Val Phe Gln Ser Val Ser Tyr Pro Gly Trp Phe Ile Ala
        115                 120                 125

acc tcc acc aca tca gga cag ccc atc ttt ctc acc aag gag aga ggc   732
Thr Ser Thr Thr Ser Gly Gln Pro Ile Phe Leu Thr Lys Glu Arg Gly
    130                 135                 140

ata act aat aac act aac ttc tac tta gat tct gtg gaa taa           774
Ile Thr Asn Asn Thr Asn Phe Tyr Leu Asp Ser Val Glu
145                 150                 155

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ctcgaatgtc actatcccaa gatgaaacct aaaaatcact ttccattctt tcttgatctt 1014

accccaccat ccactcagct gccatgccca gtttagttaa ccccccaaat gctgcttcat 1074

gcaaccttcc attcctattc cttttgccaa cccatgatgt agagatgtgg attcatgaca 1134

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 65                  70                  75                  80

Lys Pro Thr Leu Gln Leu Lys Glu Lys Asn Ile Met Asp Leu Tyr Val
                 85                  90                  95

Glu Lys Lys Ala Gln Lys Pro Phe Leu Phe Phe His Asn Lys Glu Gly
            100                 105                 110

Ser Thr Ser Val Phe Gln Ser Val Ser Tyr Pro Gly Trp Phe Ile Ala
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Pro Leu Ser Arg Ser Ile Lys Pro Val Thr Leu His Leu Ile Ala Cys
        35                  40                  45

Arg Asp Thr Glu Phe Ser Asp Lys Glu Lys Gly Asn Met Val Tyr Leu
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Gly Ile Lys Gly Lys Asp Leu Cys Leu Phe Cys Ala Glu Ile Gln Gly
 65                  70                  75                  80

Lys Pro Thr Leu Gln Leu Lys Leu Gln Gly Ser Gln Asp Asn Ile Gly
                 85                  90                  95

Lys Asp Thr Cys Trp Lys Leu Val Gly Ile His Thr Cys Ile Asn Leu
            100                 105                 110

Asp Val Arg Glu Ser Cys Phe Met Gly Thr Leu Asp Gln Trp Gly Ile
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Gly Val Gly Arg Lys Lys Trp Lys Ser Ser Phe Gln His His His Leu
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35 40 45

Cys Arg His Val Glu Thr Leu Glu Lys Asp Arg Gly Asn Pro Ile Tyr
50 55 60

Leu Gly Leu Asn Gly Leu Asn Leu Cys Leu Met Cys Ala Lys Val Gly
65 70 75 80

Asp Gln Pro Thr Leu Gln Leu Lys Glu Lys Asp Ile Met Asp Leu Tyr
85 90 95

Asn Gln Pro Glu Pro Val Lys Ser Phe Leu Phe Tyr His Ser Gln Ser
100 105 110

Gly Arg Asn Ser Thr Phe Glu Ser Val Ala Phe Pro Gly Trp Phe Ile
115 120 125

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Gly Lys Ala Asn Thr Thr Asp Phe Gly Leu Thr Met Leu Phe
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35 40 45

Tyr Leu Arg Asn Asn Gln Leu Val Ala Gly Tyr Leu Gln Gly Pro Asn

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Val Asn Leu Glu Glu Lys Ile Asp Val Val Pro Ile Glu Pro His Ala
65 70 75 80

Leu Phe Leu Gly Ile His Gly Gly Lys Met Cys Leu Ser Cys Val Lys
85 90 95

Ser Gly Asp Glu Thr Arg Leu Gln Leu Glu Ala Val Asn Ile Thr Asp
100 105 110

Leu Ser Glu Asn Arg Lys Gln Asp Lys Arg Phe Ala Phe Ile Arg Ser
115 120 125

Asp Ser Gly Pro Thr Thr Ser Phe Glu Ser Ala Ala Cys Pro Gly Trp
130 135 140

Phe Leu Cys Thr Ala Met Glu Ala Asp Gln Pro Val Ser Leu Thr Asn
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35 40 45

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Lys Asn Leu Tyr Leu Ser Cys Val Leu Lys Asp Asp Lys Pro Thr Leu
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Gln Leu Glu Ser Val Asp Pro Lys Asn Tyr Pro Lys Lys Lys Met Glu
85 90 95

Lys Arg Phe Val Phe Asn Lys Ile Glu Ile Asn Asn Lys Leu Glu Phe
100 105 110

Glu Ser Ala Gln Phe Pro Asn Trp Tyr Ile Ser Thr Ser Gln Ala Glu
115 120 125

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